RAW SEQUENCE LISTING

The Biotechnology Systems Branch of the Scientific and Technical Information Center (STIC) no errors detected.

Application Serial Number: _10/809,075_
Source: _1FWO_
Date Processed by STIC: _3/9/05_

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IFWO

RAW SEQUENCE LISTING   DATE: 03/09/2005
PATENT APPLICATION: US/10/809,075   TIME: 10:07:49

Input Set : A:\U0150863.ST25.txt
Output Set: N:\CRF4\03092005\J809075.raw

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       3 <110> APPLICANT: Dai, Ken-Shwo
       5 <120> TITLE OF INVENTION: HUMAN RPS6KA6-RELATED GENE VARIANT ASSOCIATED WITH LUNG
CANCERS
       7 <130> FILE REFERENCE: U 015086-3
C-->   9 <140> CURRENT APPLICATION NUMBER: US/10/809,075
C-->  10 <141> CURRENT FILING DATE: 2004-03-25
      12 <160> NUMBER OF SEQ ID NOS: 8
      14 <170> SOFTWARE: PatentIn version 3.1
      16 <210> SEQ ID NO: 1
      17 <211> LENGTH: 2403
      18 <212> TYPE: DNA
      19 <213> ORGANISM: Homo sapiens
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      23 <222> LOCATION: (6)..(2099)
      24 <223> OTHER INFORMATION:
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      32 Glu Val Phe Ser Gly Gly Gly Ala Ser Ser Gly Glu Val Asn Gly Leu
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      35 aaa atg gtt gat gag cca atg gaa gag gga gaa gca gat tct tgt cat     146
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      37              35                  40                  45
      39 gat gaa gga gtt gtt aaa gaa atc cct att act cat cat gtt aag gaa     194
      40 Asp Glu Gly Val Val Lys Glu Ile Pro Ile Thr His His Val Lys Glu
      41          50                  55                  60
      43 ggc tat gag aaa gca gat cct gca cag ttt gag ttg ctc aag gtt ctt     242
      44 Gly Tyr Glu Lys Ala Asp Pro Ala Gln Phe Glu Leu Leu Lys Val Leu
      45      65                  70                  75
      47 ggt cag ggg tca ttt gga aag gtt ttt ctt gtt aga aag aag acc ggt     290
      48 Gly Gln Gly Ser Phe Gly Lys Val Phe Leu Val Arg Lys Lys Thr Gly
      49 80                  85                  90                  95
      51 cct gat gct ggg cag ctc tat gca atg aag gtg tta aaa aaa gcc tct     338
      52 Pro Asp Ala Gly Gln Leu Tyr Ala Met Lys Val Leu Lys Lys Ala Ser
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      55 tta aaa gtt cga gac aga gtt cgg aca aag atg gag agg gat ata ctg     386
      56 Leu Lys Val Arg Asp Arg Val Arg Thr Lys Met Glu Arg Asp Ile Leu
      57             115                 120                 125
      59 gtg gaa gta aat cat cca ttt att gtc aaa ttg cac tat gcc ttt cag     434
      60 Val Glu Val Asn His Pro Phe Ile Val Lys Leu His Tyr Ala Phe Gln
      61             130                 135                 140
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63 act gaa ggg aaa ctg tac tta ata ctg gat ttt ctc agg gga gga gat    482
64 Thr Glu Gly Lys Leu Tyr Leu Ile Leu Asp Phe Leu Arg Gly Gly Asp
65     145                 150                 155
67 gtt ttc aca aga tta tcc aaa gag gtt ctg ttt aca gag gaa gat gtg    530
68 Val Phe Thr Arg Leu Ser Lys Glu Val Leu Phe Thr Glu Glu Asp Val
69 160                 165                 170                 175
71 aaa ttc tac ctc gca gaa ctg gcc ctt gct ttg gat cat ctg cac caa    578
72 Lys Phe Tyr Leu Ala Glu Leu Ala Leu Ala Leu Asp His Leu His Gln
73                 180                 185                 190
75 tta gga att gtt tat aga gac ctg aag cca gaa aac att ttg ctt gat    626
76 Leu Gly Ile Val Tyr Arg Asp Leu Lys Pro Glu Asn Ile Leu Leu Asp
77                 195                 200                 205
79 gaa ata gga cat atc aaa tta aca gat ttt gga ctc agc aag gag tca    674
80 Glu Ile Gly His Ile Lys Leu Thr Asp Phe Gly Leu Ser Lys Glu Ser
81     210                 215                 220
83 gta gat caa gaa aag aag gct tac tca ttt tgt ggt aca gta gag tat    722
84 Val Asp Gln Glu Lys Lys Ala Tyr Ser Phe Cys Gly Thr Val Glu Tyr
85     225                 230                 235
87 atg gct cct gaa gta gta aat agg aga ggc cat tcc cag agt gct gat    770
88 Met Ala Pro Glu Val Val Asn Arg Arg Gly His Ser Gln Ser Ala Asp
89 240                 245                 250                 255
91 tgg tgg tca tat ggt gtt ctt atg ttt gaa atg ctt act ggt act ctg    818
92 Trp Trp Ser Tyr Gly Val Leu Met Phe Glu Met Leu Thr Gly Thr Leu
93                 260                 265                 270
95 cca ttt caa ggt aaa gac aga aat gag acc atg aat atg ata tta aaa    866
96 Pro Phe Gln Gly Lys Asp Arg Asn Glu Thr Met Asn Met Ile Leu Lys
97                 275                 280                 285
99 gca aaa ctt gga atg cct caa ttt ctt agt gct gaa gca caa agt ctt    914
100 Ala Lys Leu Gly Met Pro Gln Phe Leu Ser Ala Glu Ala Gln Ser Leu
101                 290                 295                 300
103 cta agg atg tta ttc aaa agg aat cca gca aat aga ttg gga tca gaa    962
104 Leu Arg Met Leu Phe Lys Arg Asn Pro Ala Asn Arg Leu Gly Ser Glu
105     305                 310                 315
107 gga gtt gaa gaa atc aaa aga cat ctg ttt ttt gca aat att gac tgg    1010
108 Gly Val Glu Glu Ile Lys Arg His Leu Phe Phe Ala Asn Ile Asp Trp
109 320                 325                 330                 335
111 gat aaa tta tat aaa aga gaa gtt caa cct cct ttc aaa cct gct tct    1058
112 Asp Lys Leu Tyr Lys Arg Glu Val Gln Pro Pro Phe Lys Pro Ala Ser
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115 gga aaa cca gat gat act ttt tgt ttt gat cct gaa ttt act gca aaa    1106
116 Gly Lys Pro Asp Asp Thr Phe Cys Phe Asp Pro Glu Phe Thr Ala Lys
117                 355                 360                 365
119 aca cct aaa gat tct ccc ggt ttg cca gcc agt gca aat gct cat cag    1154
120 Thr Pro Lys Asp Ser Pro Gly Leu Pro Ala Ser Ala Asn Ala His Gln
121                 370                 375                 380
123 ctc ttc aaa gga ttc agc ttt gtt gca act tct att gca gaa gaa tat    1202
124 Leu Phe Lys Gly Phe Ser Phe Val Ala Thr Ser Ile Ala Glu Glu Tyr
125     385                 390                 395
127 aaa atc act cct atc aca agt gca aat gta tta cca att gtt cag ata    1250
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128 Lys Ile Thr Pro Ile Thr Ser Ala Asn Val Leu Pro Ile Val Gln Ile
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131 aat gga aat gct gca caa ttt ggt gaa gta tat gaa ttg aag gag gat      1298
132 Asn Gly Asn Ala Ala Gln Phe Gly Glu Val Tyr Glu Leu Lys Glu Asp
133                 420                 425                 430
135 att ggt gtt ggc tcc tac tct gtt tgc aag cga tgc ata cat gca act      1346
136 Ile Gly Val Gly Ser Tyr Ser Val Cys Lys Arg Cys Ile His Ala Thr
137                 435                 440                 445
139 acc aac atg gaa ttt gca gtg aag atc att gac aaa agt aag cga gac      1394
140 Thr Asn Met Glu Phe Ala Val Lys Ile Ile Asp Lys Ser Lys Arg Asp
141             450                 455                 460
143 cct tca gaa gag att gaa ata ttg atg cgc tat gga caa cat ccc aac      1442
144 Pro Ser Glu Glu Ile Glu Ile Leu Met Arg Tyr Gly Gln His Pro Asn
145         465                 470                 475
147 att att act ttg aag gat gtc ttt gat gat ggt aga tat gtt tac ctt      1490
148 Ile Ile Thr Leu Lys Asp Val Phe Asp Asp Gly Arg Tyr Val Tyr Leu
149 480                 485                 490                 495
151 gtt acg gat tta atg aaa gga gga gag tta ctt gac cgt att ctc aaa      1538
152 Val Thr Asp Leu Met Lys Gly Gly Glu Leu Leu Asp Arg Ile Leu Lys
153             500                 505                 510
155 caa aaa tgt ttc tcg gaa cgg gag gct agt gat ata cta tat gta ata      1586
156 Gln Lys Cys Phe Ser Glu Arg Glu Ala Ser Asp Ile Leu Tyr Val Ile
157             515                 520                 525
159 agt aag aca gtt gac tat ctt cat tgt caa gga gtt gtt cat cgt gat      1634
160 Ser Lys Thr Val Asp Tyr Leu His Cys Gln Gly Val Val His Arg Asp
161             530                 535                 540
163 ctt aaa cct agt aat att tta tac atg gat gaa tca gcc agt gca gat      1682
164 Leu Lys Pro Ser Asn Ile Leu Tyr Met Asp Glu Ser Ala Ser Ala Asp
165         545                 550                 555
167 tca atc agg ata tgt gat ttt ggg ttt gca aaa caa ctt cga gga gaa      1730
168 Ser Ile Arg Ile Cys Asp Phe Gly Phe Ala Lys Gln Leu Arg Gly Glu
169 560                 565                 570                 575
171 aat gga ctt ctc tta act cca tgc tac act gca aac ttt gtt gca cct      1778
172 Asn Gly Leu Leu Leu Thr Pro Cys Tyr Thr Ala Asn Phe Val Ala Pro
173                 580                 585                 590
175 gag gtt ctt atg caa cag gga tat gat gct gct tgt gat atc tgg agt      1826
176 Glu Val Leu Met Gln Gln Gly Tyr Asp Ala Ala Cys Asp Ile Trp Ser
177             595                 600                 605
179 tta gga gtc ctt ttt tac aca atg ttg gct ggc tac act cca ttt gct      1874
180 Leu Gly Val Leu Phe Tyr Thr Met Leu Ala Gly Tyr Thr Pro Phe Ala
181             610                 615                 620
183 aat ggc ccc aat gat act cct gaa gag ata ctg ctg cgt ata ggc aat      1922
184 Asn Gly Pro Asn Asp Thr Pro Glu Glu Ile Leu Leu Arg Ile Gly Asn
185         625                 630                 635
187 gga aaa ttc tct ttg agt ggt gga aac tgg gac aat att tca gac gga      1970
188 Gly Lys Phe Ser Leu Ser Gly Gly Asn Trp Asp Asn Ile Ser Asp Gly
189 640                 645                 650                 655
191 gca aag gga gca atg gtt gca aca tac tct gcc ctg act cac aag acc      2018
192 Ala Lys Gly Ala Met Val Ala Thr Tyr Ser Ala Leu Thr His Lys Thr
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195 ttt caa cca gtc cta gag cct gta gct gct tca agc tta gcc cag cga      2066
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197               675             680             685
199 cgg agc atg aaa aag cga aca tca act ggc ctg taagatttgt ggtgttccta    2119
200 Arg Ser Met Lys Lys Arg Thr Ser Thr Gly Leu
201           690             695
203 ggccaaactg gatgaagatg aaattaaatg tgtggctttt ttcctattct tatcaaaggc    2179
205 atcgttgtct gctaaattac ttgaatatta agtaatatta aatccccatt tttaggggaa    2239
207 gtgagattta aaaaaccatt cacaggtcca caatattcat actatgtgtt tgcagtagtg    2299
209 ttcaagtgtt tatttaagca tataattggt gtccaccagg tcctcacaac ttctctgcac    2359
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216 <212> TYPE: PRT
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230         35                  40                  45
233 Glu Gly Val Val Lys Glu Ile Pro Ile Thr His His Val Lys Glu Gly
234     50                  55                  60
237 Tyr Glu Lys Ala Asp Pro Ala Gln Phe Glu Leu Leu Lys Val Leu Gly
238 65                  70                  75                  80
241 Gln Gly Ser Phe Gly Lys Val Phe Leu Val Arg Lys Lys Thr Gly Pro
242               85                  90                  95
245 Asp Ala Gly Gln Leu Tyr Ala Met Lys Val Leu Lys Lys Ala Ser Leu
246               100                 105                 110
249 Lys Val Arg Asp Arg Val Arg Thr Lys Met Glu Arg Asp Ile Leu Val
250           115                 120                 125
253 Glu Val Asn His Pro Phe Ile Val Lys Leu His Tyr Ala Phe Gln Thr
254     130                 135                 140
257 Glu Gly Lys Leu Tyr Leu Ile Leu Asp Phe Leu Arg Gly Gly Asp Val
258 145                 150                 155                 160
261 Phe Thr Arg Leu Ser Lys Glu Val Leu Phe Thr Glu Glu Asp Val Lys
262               165                 170                 175
265 Phe Tyr Leu Ala Glu Leu Ala Leu Ala Leu Asp His Leu His Gln Leu
266               180                 185                 190
269 Gly Ile Val Tyr Arg Asp Leu Lys Pro Glu Asn Ile Leu Leu Asp Glu
270           195                 200                 205
273 Ile Gly His Ile Lys Leu Thr Asp Phe Gly Leu Ser Lys Glu Ser Val
274     210                 215                 220
277 Asp Gln Glu Lys Lys Ala Tyr Ser Phe Cys Gly Thr Val Glu Tyr Met
278 225                 230                 235                 240
281 Ala Pro Glu Val Val Asn Arg Arg Gly His Ser Gln Ser Ala Asp Trp
282               245                 250                 255
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285 Trp Ser Tyr Gly Val Leu Met Phe Glu Met Leu Thr Gly Thr Leu Pro
286             260                 265             270
289 Phe Gln Gly Lys Asp Arg Asn Glu Thr Met Asn Met Ile Leu Lys Ala
290         275             280             285
293 Lys Leu Gly Met Pro Gln Phe Leu Ser Ala Glu Ala Gln Ser Leu Leu
294     290             295             300
297 Arg Met Leu Phe Lys Arg Asn Pro Ala Asn Arg Leu Gly Ser Glu Gly
298 305             310             315                 320
301 Val Glu Glu Ile Lys Arg His Leu Phe Phe Ala Asn Ile Asp Trp Asp
302             325             330             335
305 Lys Leu Tyr Lys Arg Glu Val Gln Pro Pro Phe Lys Pro Ala Ser Gly
306             340             345             350
309 Lys Pro Asp Asp Thr Phe Cys Phe Asp Pro Glu Phe Thr Ala Lys Thr
310         355             360             365
313 Pro Lys Asp Ser Pro Gly Leu Pro Ala Ser Ala Asn Ala His Gln Leu
314     370             375             380
317 Phe Lys Gly Phe Ser Phe Val Ala Thr Ser Ile Ala Glu Glu Tyr Lys
318 385             390             395                 400
321 Ile Thr Pro Ile Thr Ser Ala Asn Val Leu Pro Ile Val Gln Ile Asn
322             405             410             415
325 Gly Asn Ala Ala Gln Phe Gly Glu Val Tyr Glu Leu Lys Glu Asp Ile
326             420             425             430
329 Gly Val Gly Ser Tyr Ser Val Cys Lys Arg Cys Ile His Ala Thr Thr
330         435             440             445
333 Asn Met Glu Phe Ala Val Lys Ile Ile Asp Lys Ser Lys Arg Asp Pro
334     450             455             460
337 Ser Glu Glu Ile Glu Ile Leu Met Arg Tyr Gly Gln His Pro Asn Ile
338 465             470             475                 480
341 Ile Thr Leu Lys Asp Val Phe Asp Asp Gly Arg Tyr Val Tyr Leu Val
342             485             490             495
345 Thr Asp Leu Met Lys Gly Gly Glu Leu Leu Asp Arg Ile Leu Lys Gln
346             500             505             510
349 Lys Cys Phe Ser Glu Arg Glu Ala Ser Asp Ile Leu Tyr Val Ile Ser
350         515             520             525
353 Lys Thr Val Asp Tyr Leu His Cys Gln Gly Val Val His Arg Asp Leu
354     530             535             540
357 Lys Pro Ser Asn Ile Leu Tyr Met Asp Glu Ser Ala Ser Ala Asp Ser
358 545             550             555                 560
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362             565             570             575
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366             580             585             590
369 Val Leu Met Gln Gln Gly Tyr Asp Ala Ala Cys Asp Ile Trp Ser Leu
370         595             600             605
373 Gly Val Leu Phe Tyr Thr Met Leu Ala Gly Tyr Thr Pro Phe Ala Asn
374     610             615             620
377 Gly Pro Asn Asp Thr Pro Glu Glu Ile Leu Leu Arg Ile Gly Asn Gly
378 625             630             635             640
381 Lys Phe Ser Leu Ser Gly Gly Asn Trp Asp Asn Ile Ser Asp Gly Ala
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VERIFICATION SUMMARY   DATE: 03/09/2005
PATENT APPLICATION: **US/10/809,075**   TIME: 10:07:50

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L:9 M:270 C: Current Application Number differs, Replaced Current Application Number
L:10 M:271 C: Current Filing Date differs, Replaced Current Filing Date
L:26 M:258 W: Mandatory Feature missing, <223> Blank for SEQ#:1,Line#:24
L:407 M:258 W: Mandatory Feature missing, <223> Blank for SEQ#:3,Line#:405